

**In the Claims:**

Please amend claims 24 and 33 as follows:

24. (Amended) A method for identifying a compound capable of interfering with binding of an TRAC1 polypeptide or fragment thereof, the method comprising the steps of:

(i) combining an TRAC1 polypeptide or fragment thereof with an E2 ubiquitin-conjugating enzyme polypeptide and the compound, wherein the TRAC1 polypeptide or fragment thereof is encoded by a nucleic acid that hybridizes under stringent conditions to a nucleic acid encoding a polypeptide having an amino acid sequence of SEQ ID NO:1; and

(ii) determining the binding of the TRAC1 polypeptide or fragment thereof to the E2 ubiquitin-conjugating enzyme polypeptide.

33. (Amended) An isolated complex comprising a TRAC1 polypeptide or fragment thereof bound to an E2 ubiquitin-conjugating enzyme polypeptide, wherein the TRAC1 polypeptide or fragment thereof is encoded by a nucleic acid that hybridizes under stringent conditions to a nucleic acid encoding a polypeptide having an amino acid sequence of SEQ ID NO:1.

**REMARKS**

Claims 1-46 are pending in this application. Claims 24 and 33 have been amended. The amendments to claims 24 and 33 correct the assigned identifiers for SEQ ID NO: designated in these claims. The claim language in claims 24 and 33 refer to amino acid sequences, whereas SEQ ID NO:2 is the corresponding nucleic acid sequence for which the amended SEQ ID NO:1 pertains.